## **Amendments to the Claims:**

- 1-27. (canceled)
- 28. (currently amended) An isolated <u>native sequence</u> polypeptide having at least 80% amino acid sequence identity to:
  - (a) the amino acid sequence of the polypeptide of SEQ ID NO:374;
- (b) the amino acid sequence of the polypeptide of SEQ ID NO:374, lacking its associated signal peptide; or
- (c) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203465;

wherein the nucleic acid encoding the polypeptide is amplified in lung or colon tumors.

- 29. (currently amended) The isolated <u>native sequence</u> polypeptide of Claim 28 having at least 85% amino acid sequence identity to:
  - (a) the amino acid sequence of the polypeptide of SEQ ID NO:374;
- (b) the amino acid sequence of the polypeptide of SEQ ID NO:374, lacking its associated signal peptide; or
- (c) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203465;

wherein the nucleic acid encoding the polypeptide is amplified in lung or colon tumors.

- 30. (currently amended) The isolated <u>native sequence</u> polypeptide of Claim 28 having at least 90% amino acid sequence identity to:
  - (a) the amino acid sequence of the polypeptide of SEQ ID NO:374;
- (b) the amino acid sequence of the polypeptide of SEQ ID NO:374, lacking its associated signal peptide; or
- (c) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203465;

wherein the nucleic acid encoding the polypeptide is amplified in lung or colon tumors.

- 31. (currently amended) The isolated <u>native sequence</u> polypeptide of Claim 28 having at least 95% amino acid sequence identity to:
  - (a) the amino acid sequence of the polypeptide of SEQ ID NO:374;
- (b) the amino acid sequence of the polypeptide of SEQ ID NO:374, lacking its associated signal peptide; or
- (c) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203465;

wherein the nucleic acid encoding the polypeptide is amplified in lung or colon tumors.

- 32. (currently amended) The isolated <u>native sequence</u> polypeptide of Claim 28 having at least 99% amino acid sequence identity to:
  - (a) the amino acid sequence of the polypeptide of SEQ ID NO:374;
- (b) the amino acid sequence of the polypeptide of SEQ ID NO:374, lacking its associated signal peptide; or
- (c) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203465;

wherein the nucleic acid encoding the polypeptide is amplified in lung or colon tumors.

- 33. (previously presented) An isolated polypeptide comprising:
- (a) the amino acid sequence of the polypeptide of SEQ ID NO:374;
- (b) the amino acid sequence of the polypeptide of SEQ ID NO:374, lacking its associated signal peptide; or
- (c) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203465.
- 34. (previously presented) The isolated polypeptide of Claim 33 comprising the amino acid sequence of the polypeptide of SEQ ID NO:374.
- 35. (previously presented) The isolated polypeptide of Claim 33 comprising the amino acid sequence of the polypeptide of SEQ ID NO:374, lacking its associated signal peptide.

- 36. (canceled)
- 37. (canceled)
- 38. (previously presented) The isolated polypeptide of Claim 33 comprising the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203465.
- 39. (previously presented) A chimeric polypeptide comprising a polypeptide according to Claim 28 fused to a heterologous polypeptide.
- 40. (previously presented) The chimeric polypeptide of Claim 39, wherein said heterologous polypeptide is an epitope tag or an Fc region of an immunoglobulin.